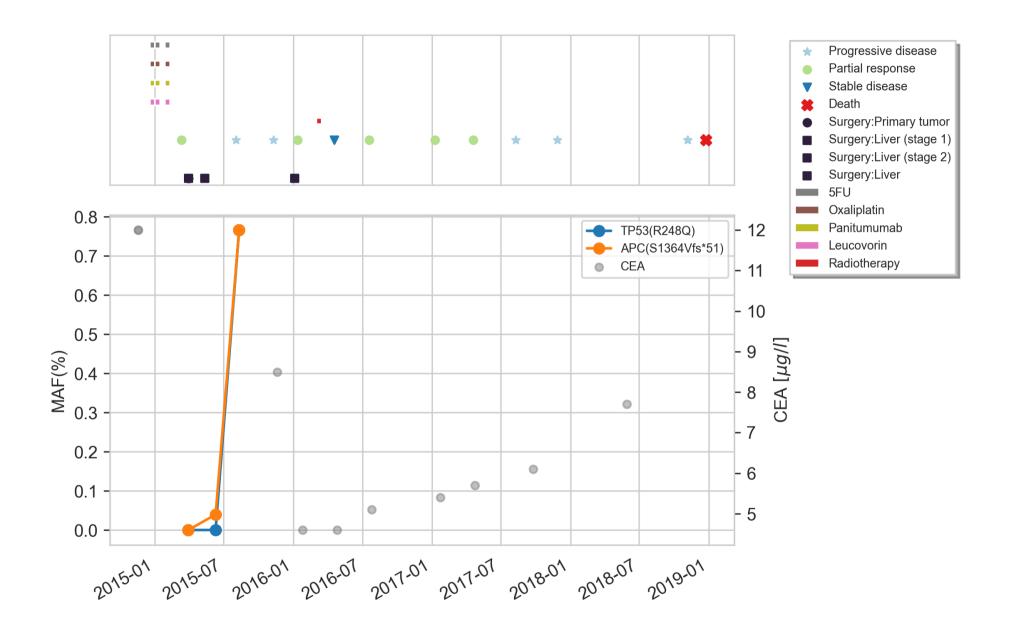
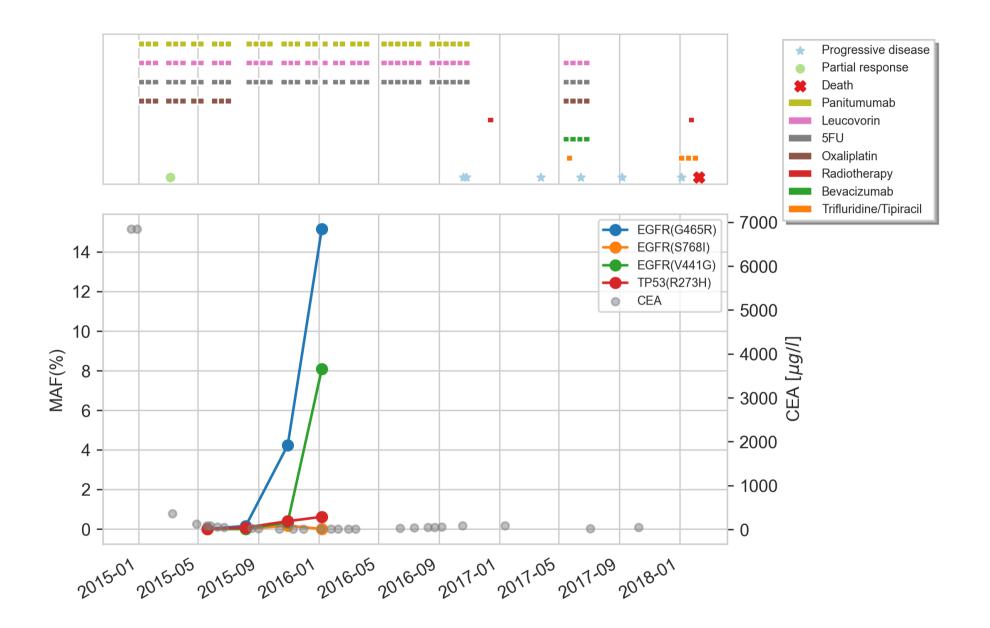
Supplementary Data 1. Individual plots of all 52 patients showing ctDNA dynamics in the bottom panel by depicting the mutant allele frequencies (MAF) of the identified somatic mutations after correction for germline and hematopoietic variants (left y-axis) as well as the CEA levels (right y-axis). Following the same x-axis, the top panel provides information about the patient's treatment, surgeries, and radiological assessments.

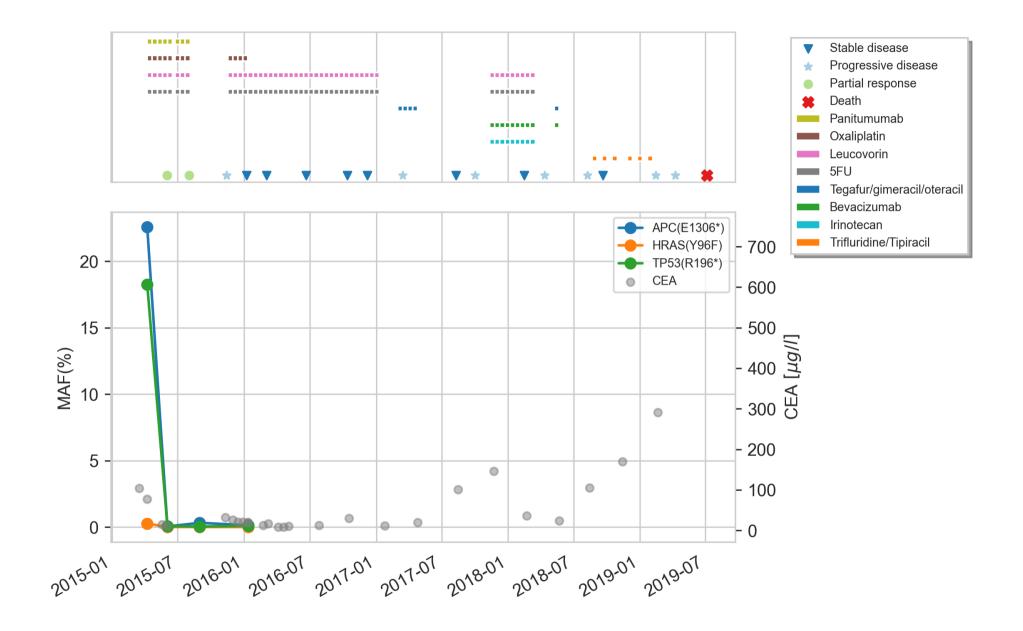
Patient 2



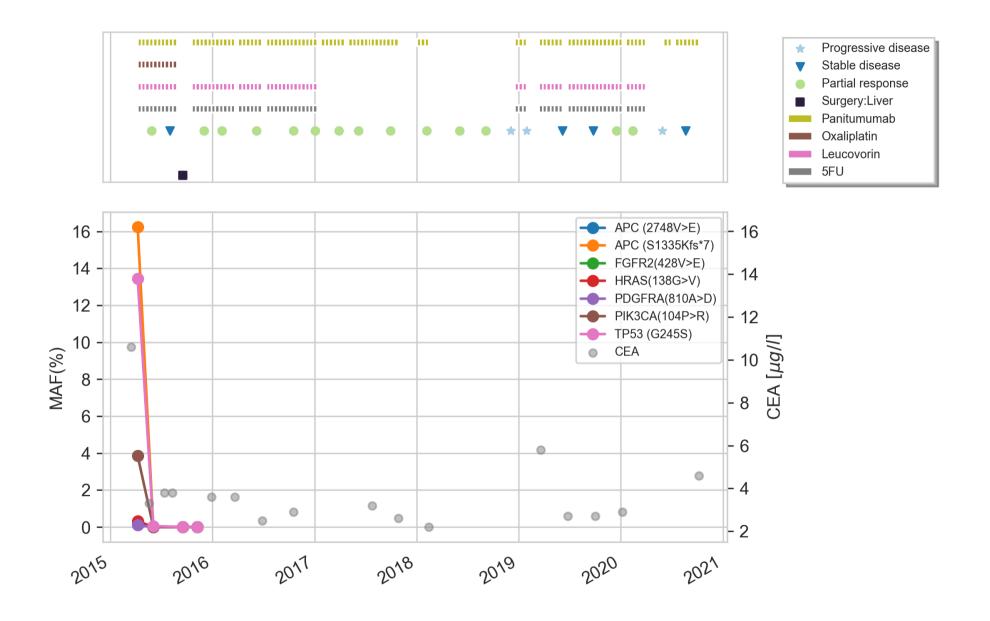
Patient 4



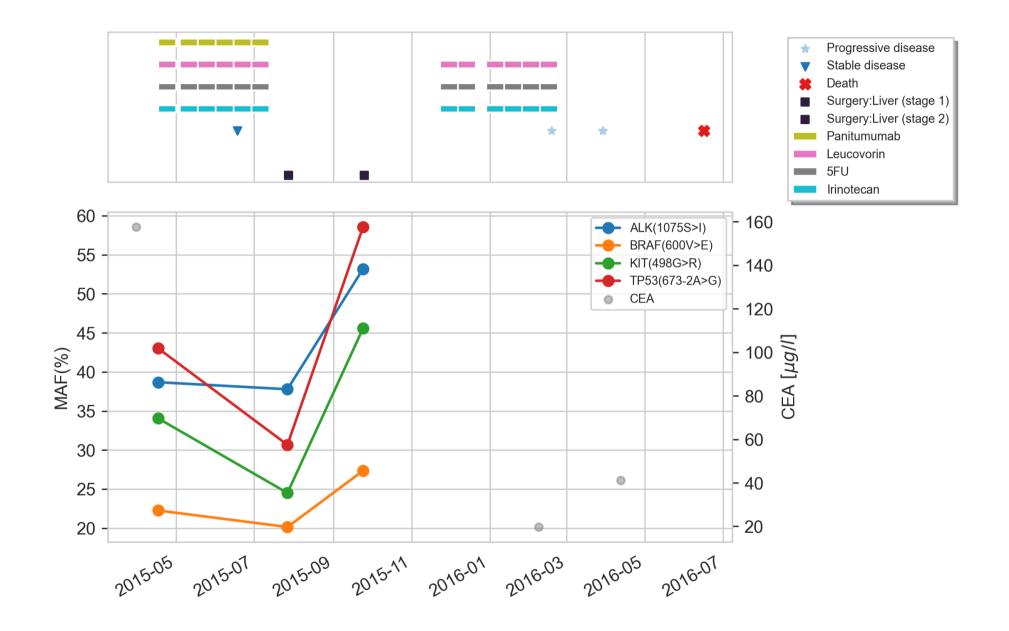
Patient 12



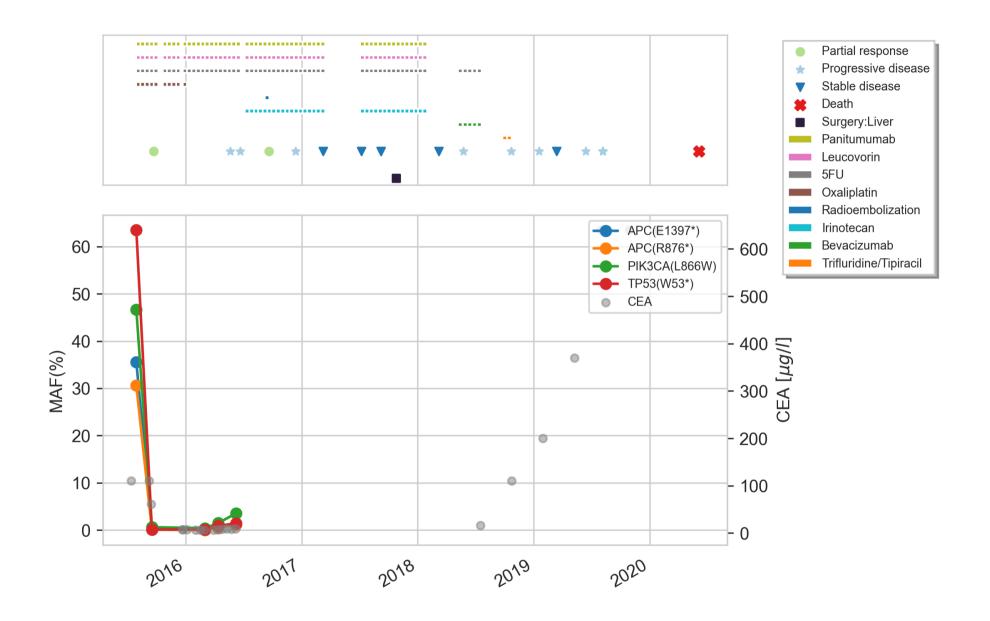
Patient 13

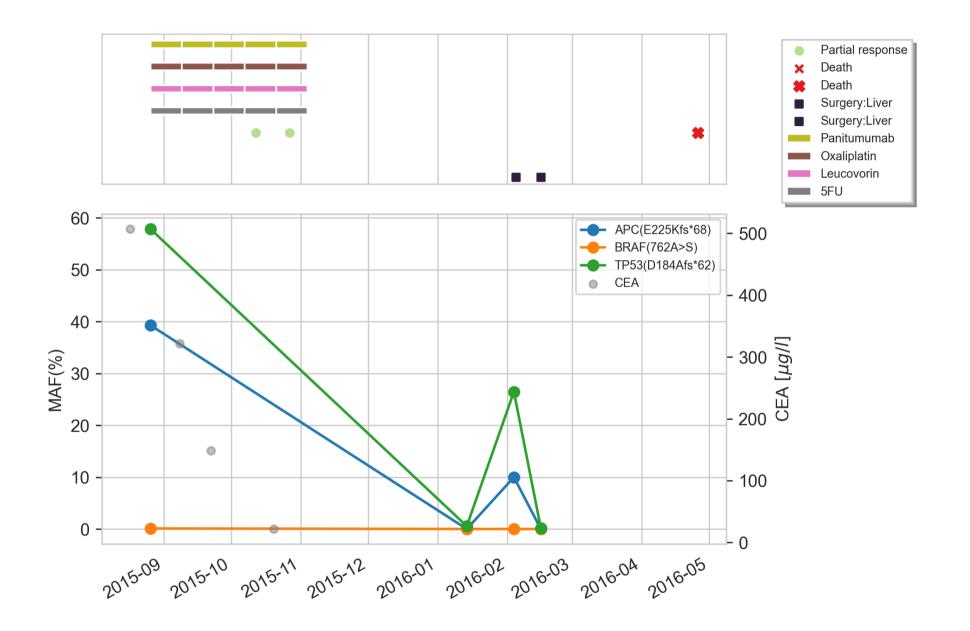


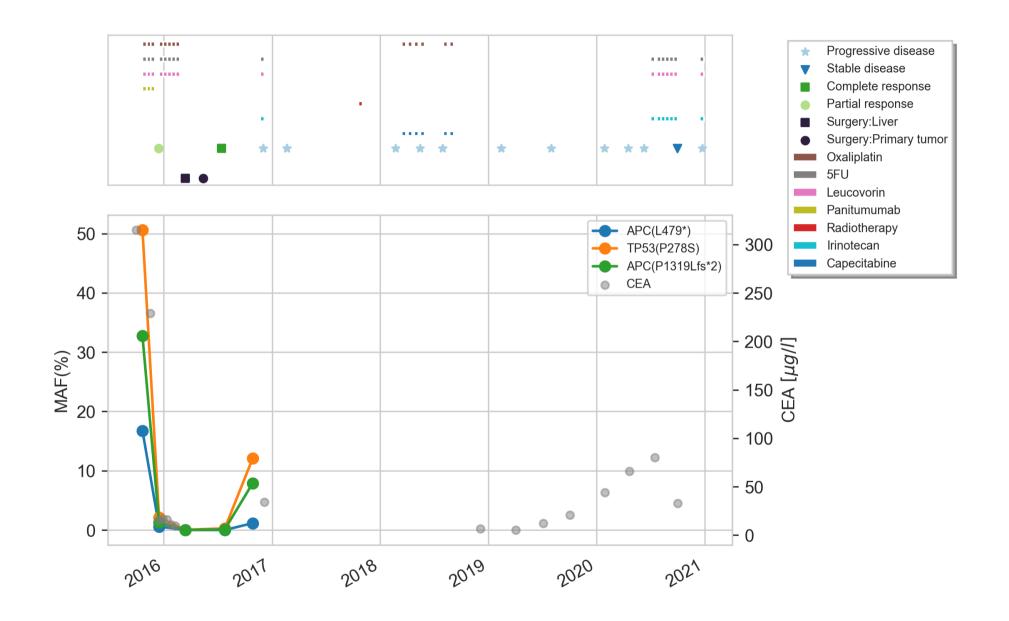
Patient 14

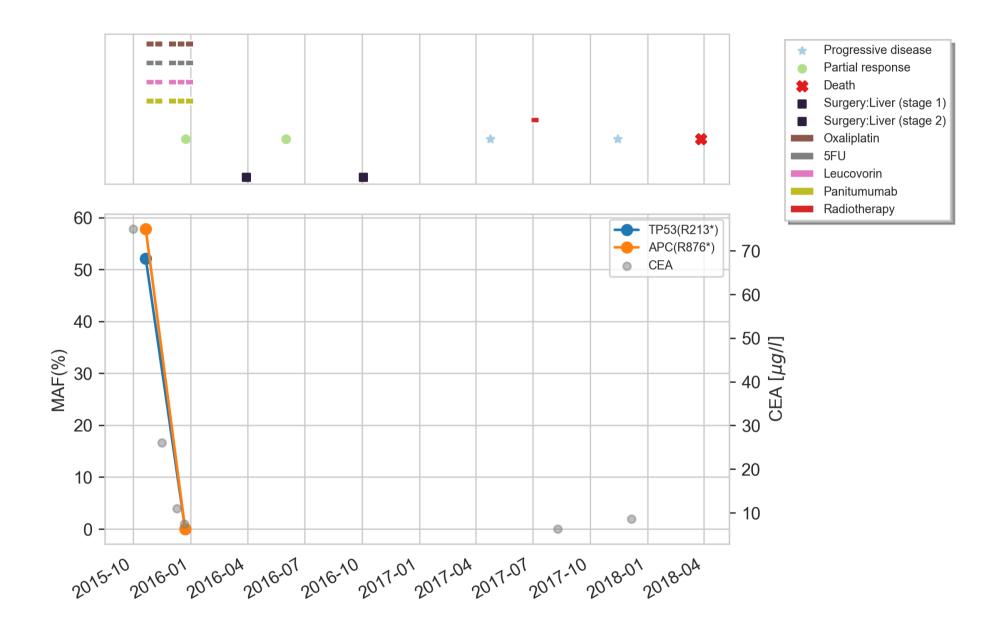


Patient 27

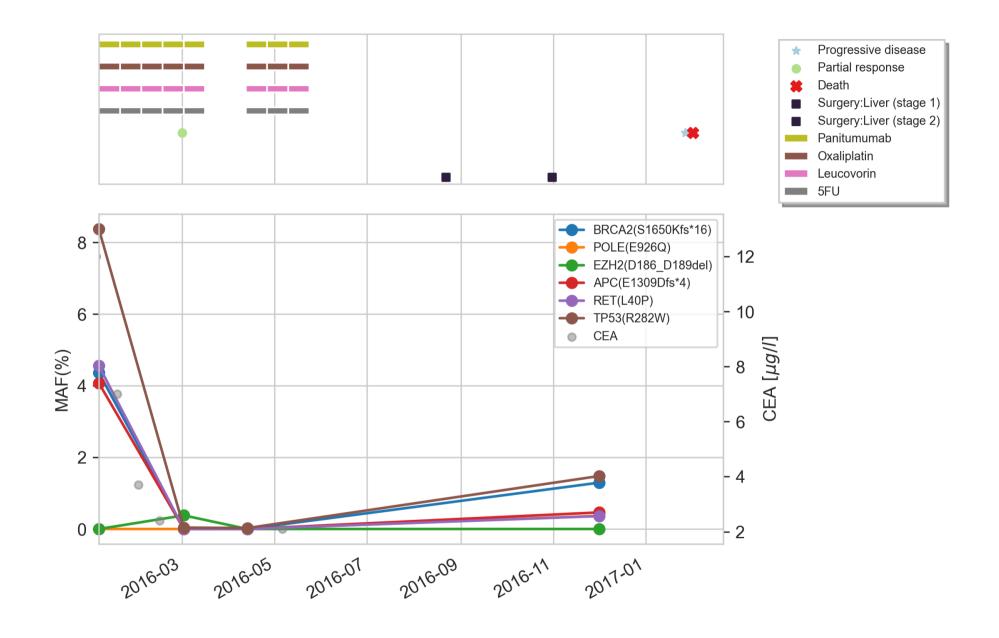


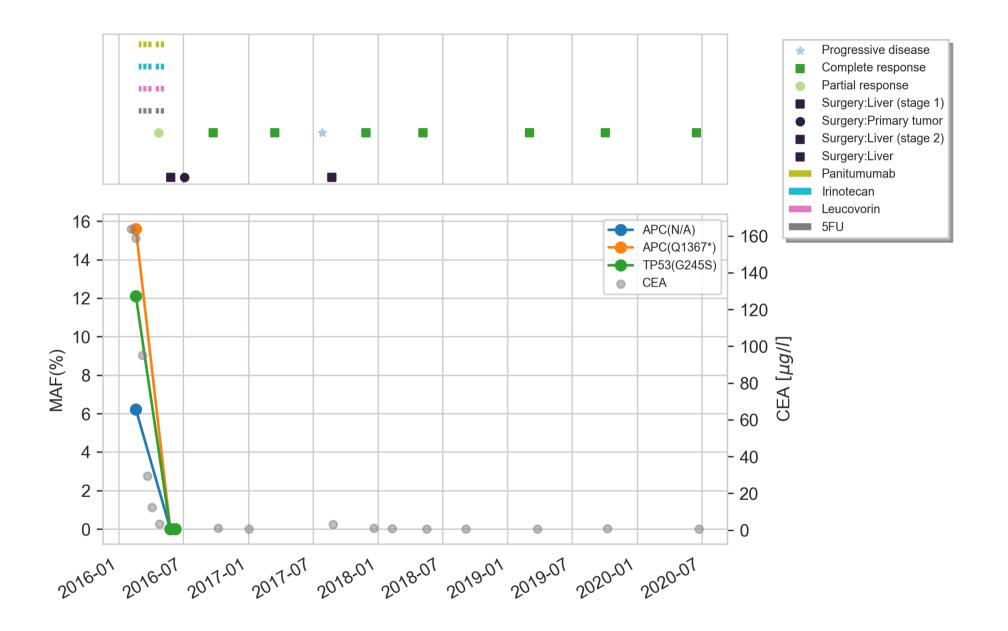


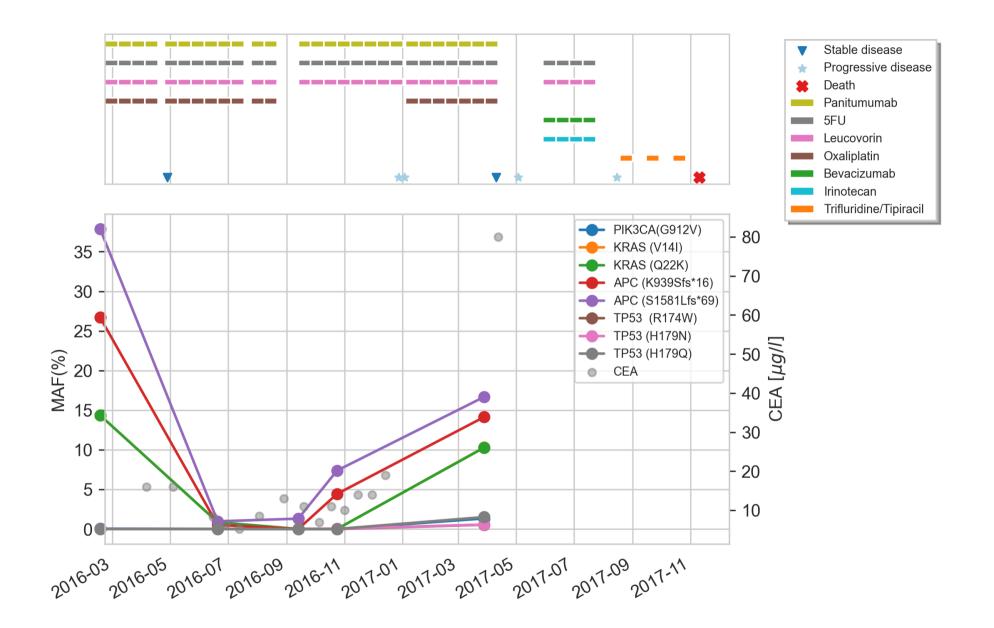


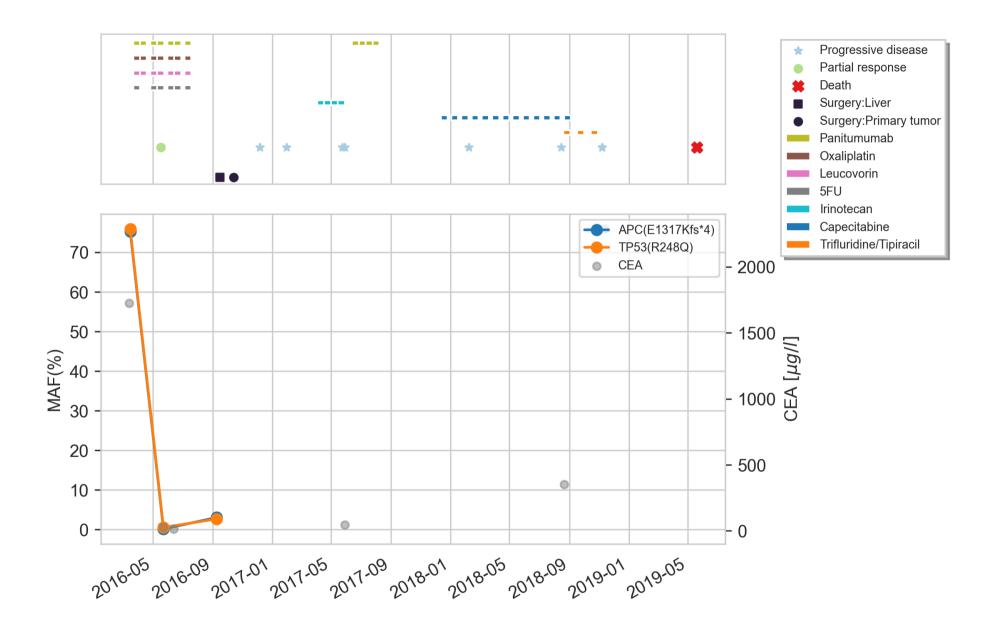


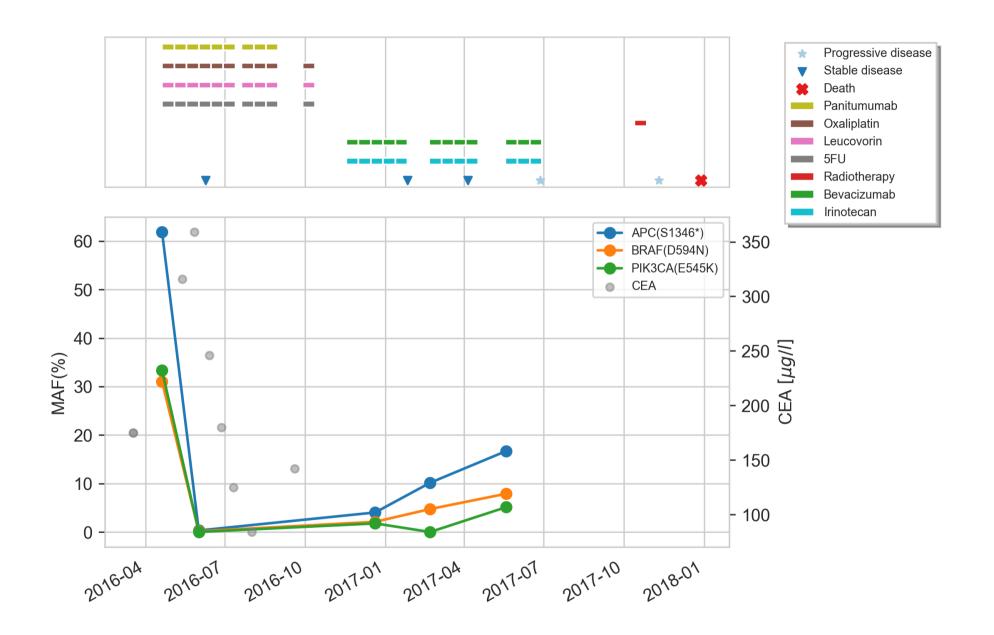
Patient 45

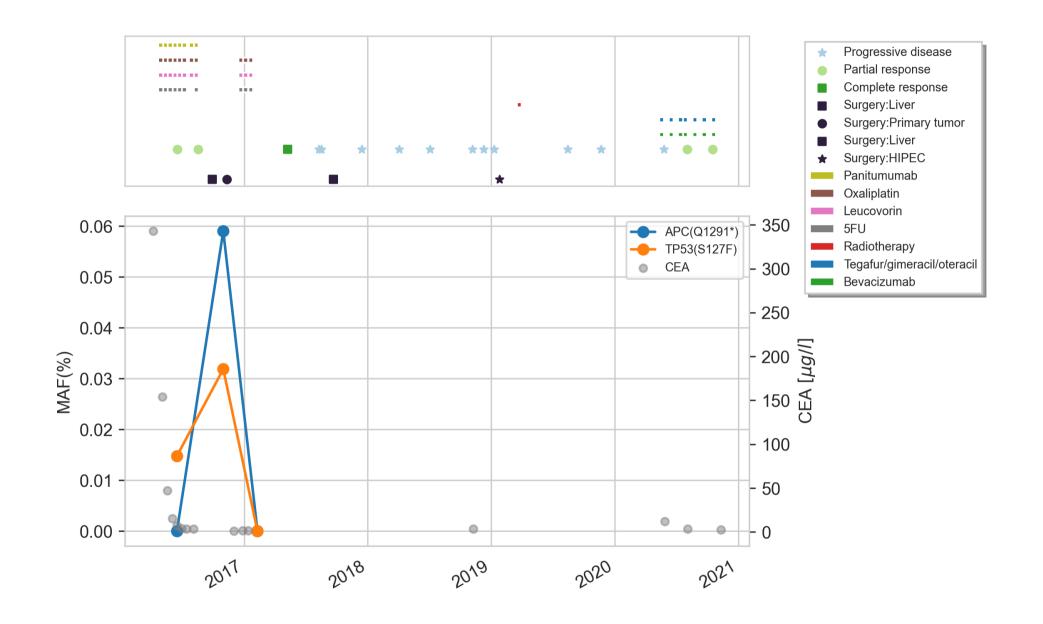


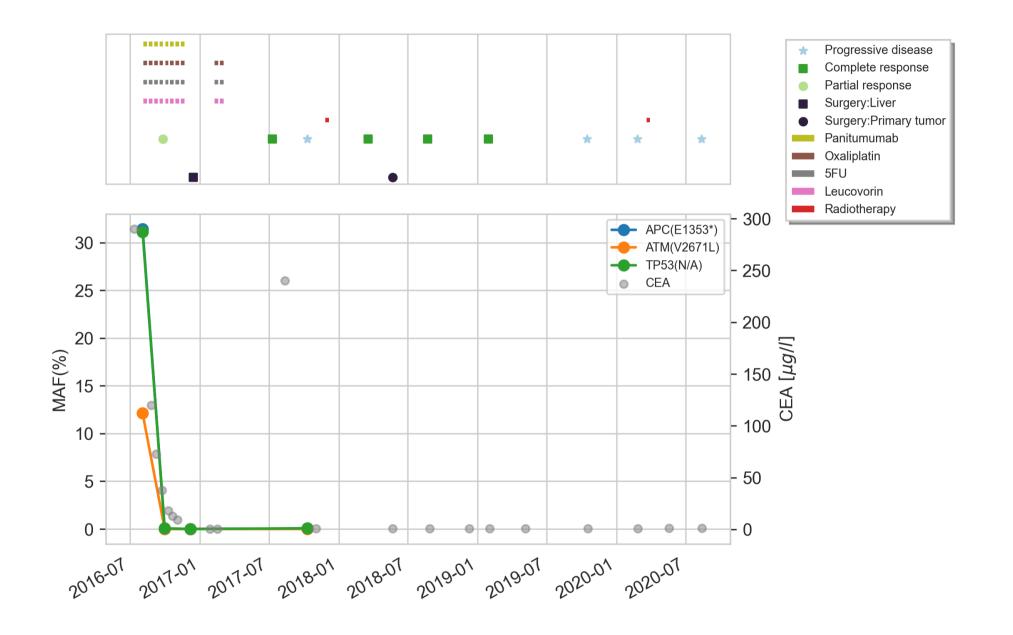


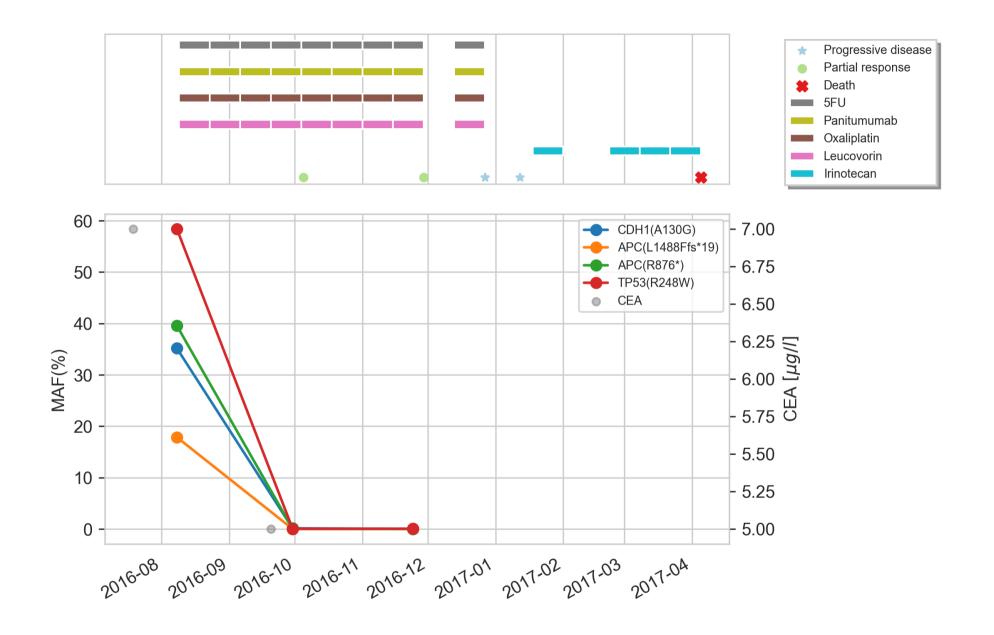




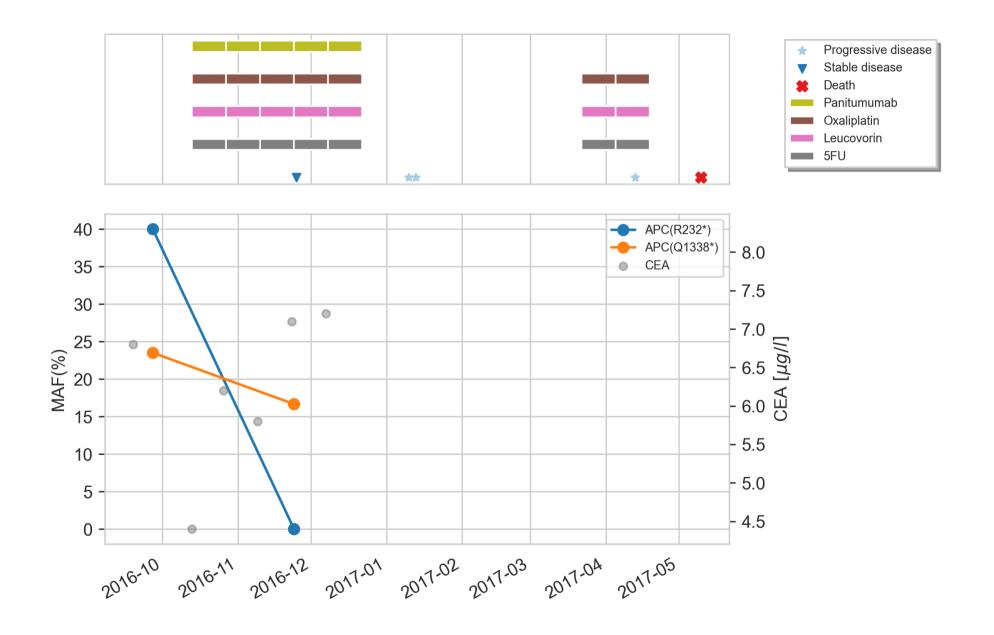


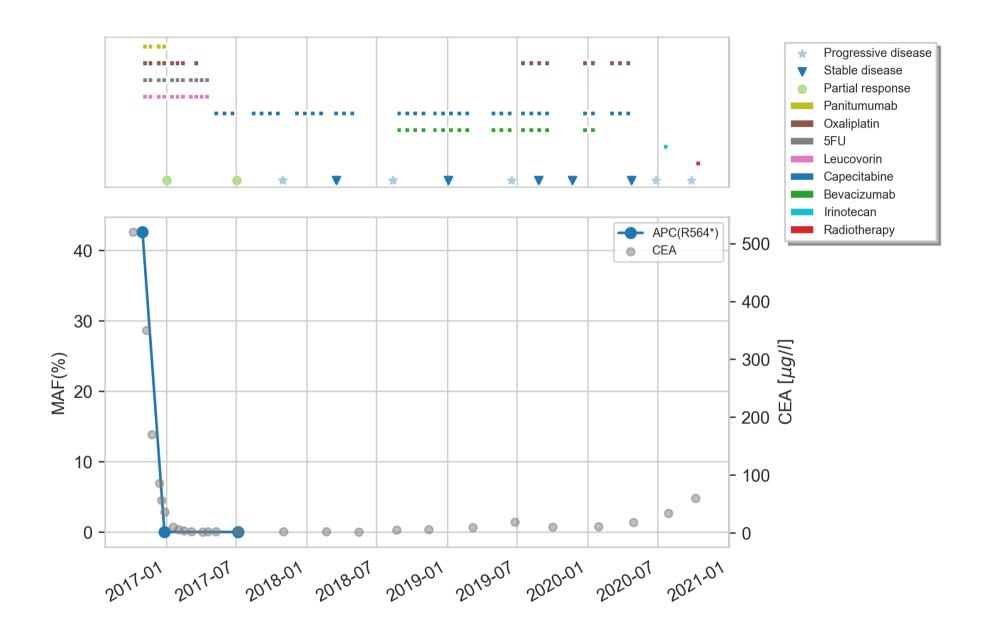


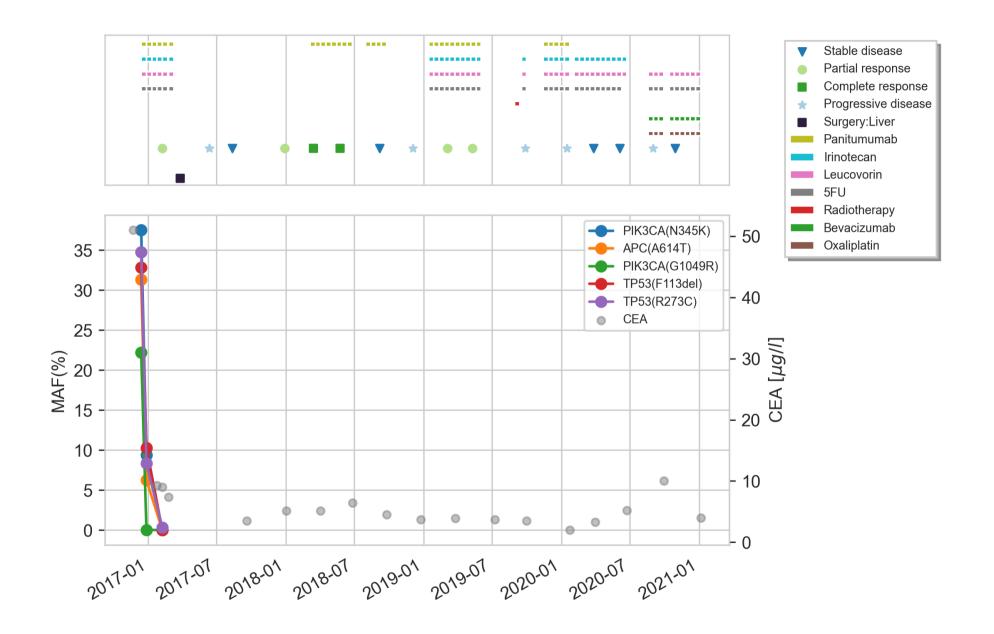




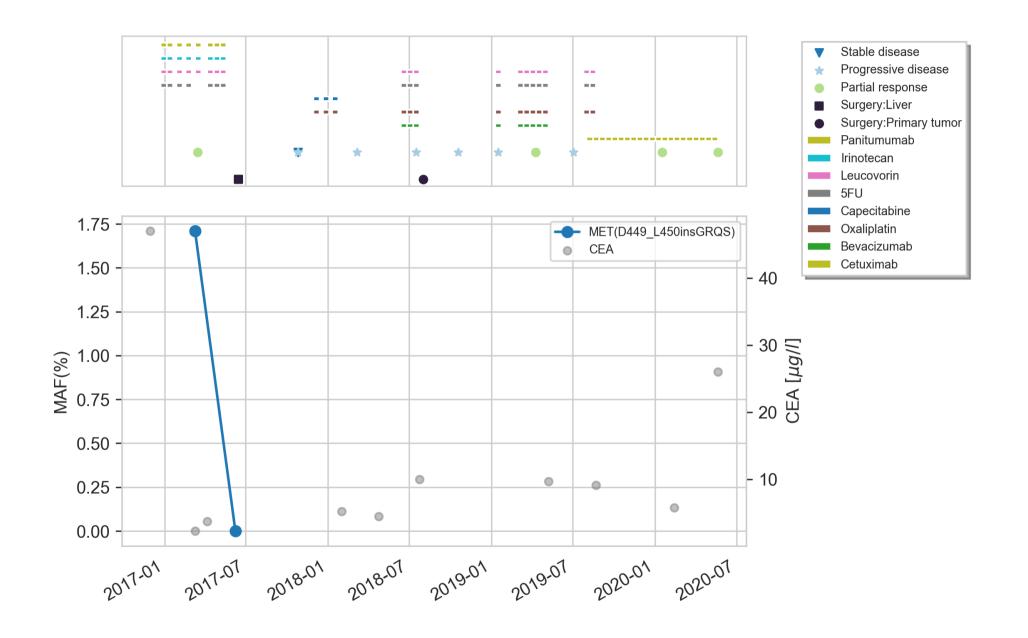
Patient 124



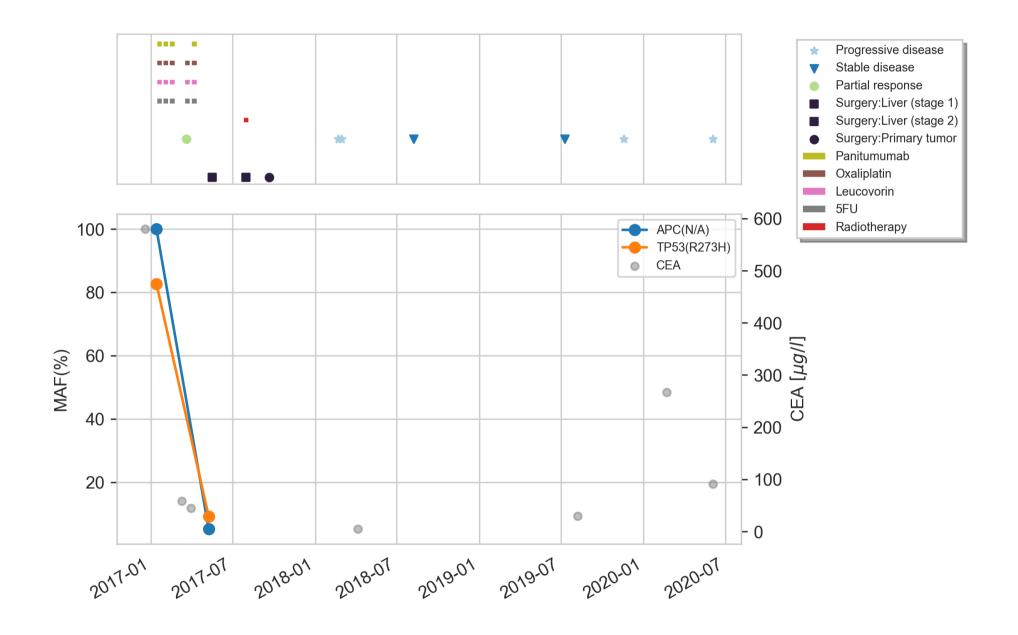




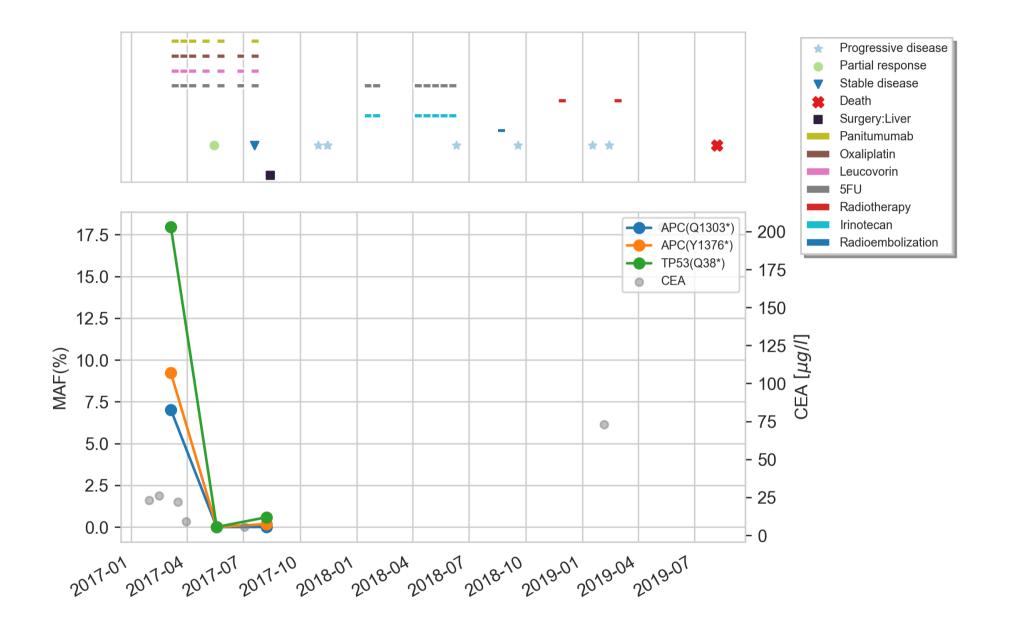
Patient 140

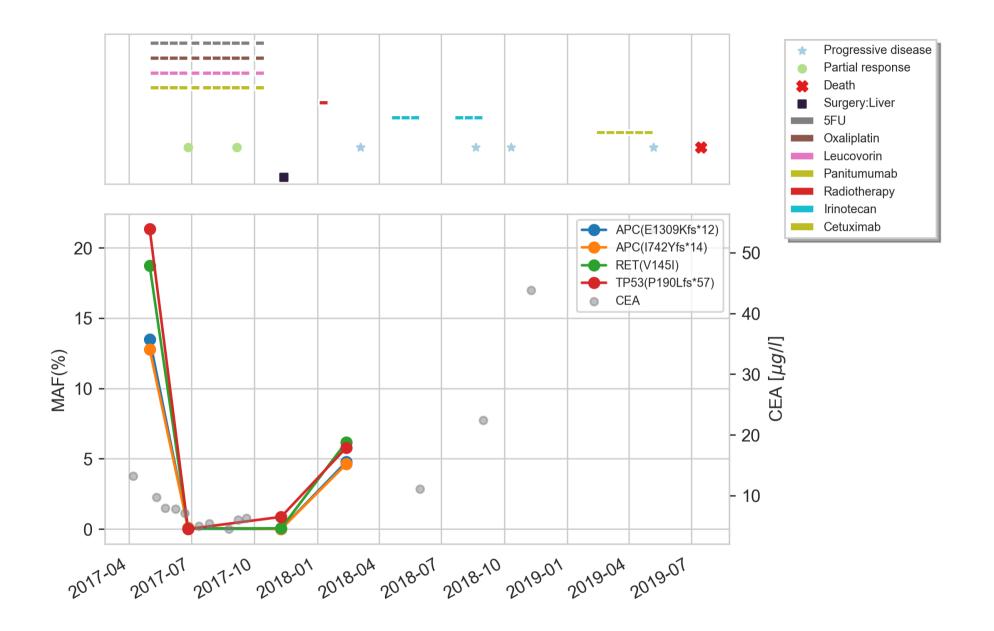


Patient 148

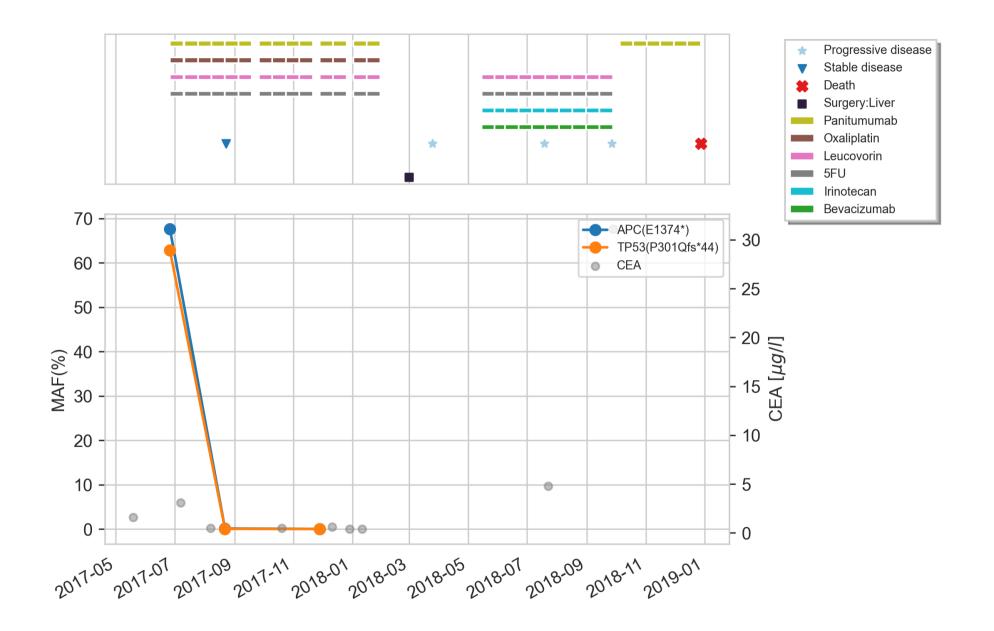


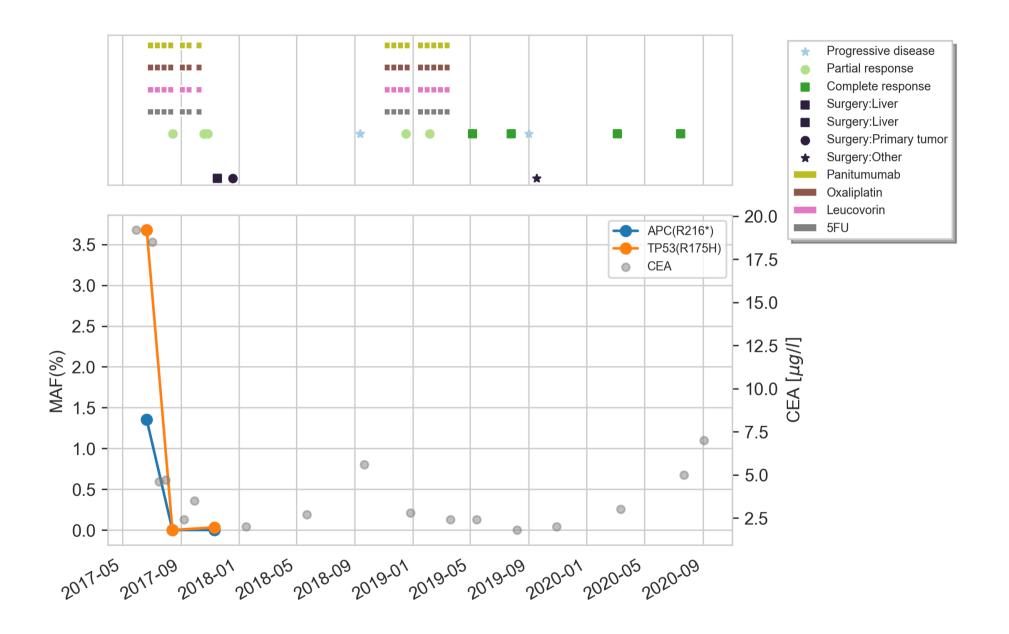
Patient 157

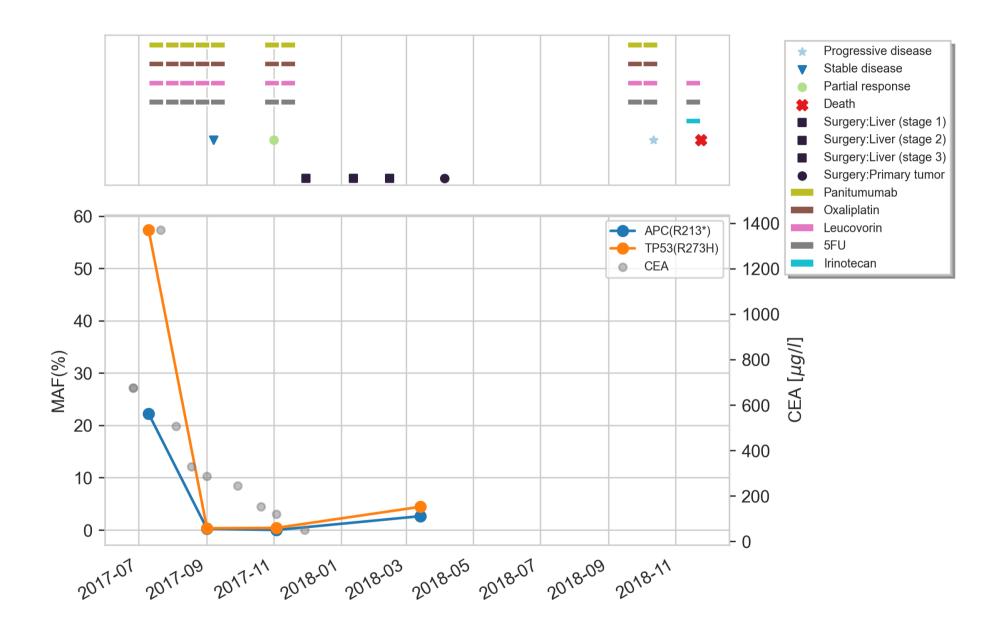


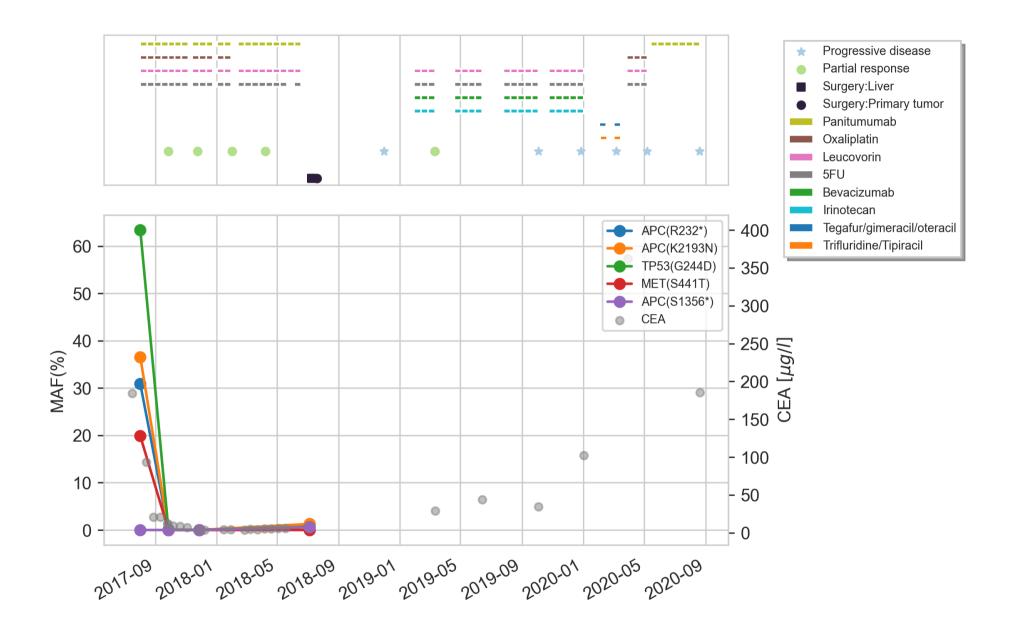


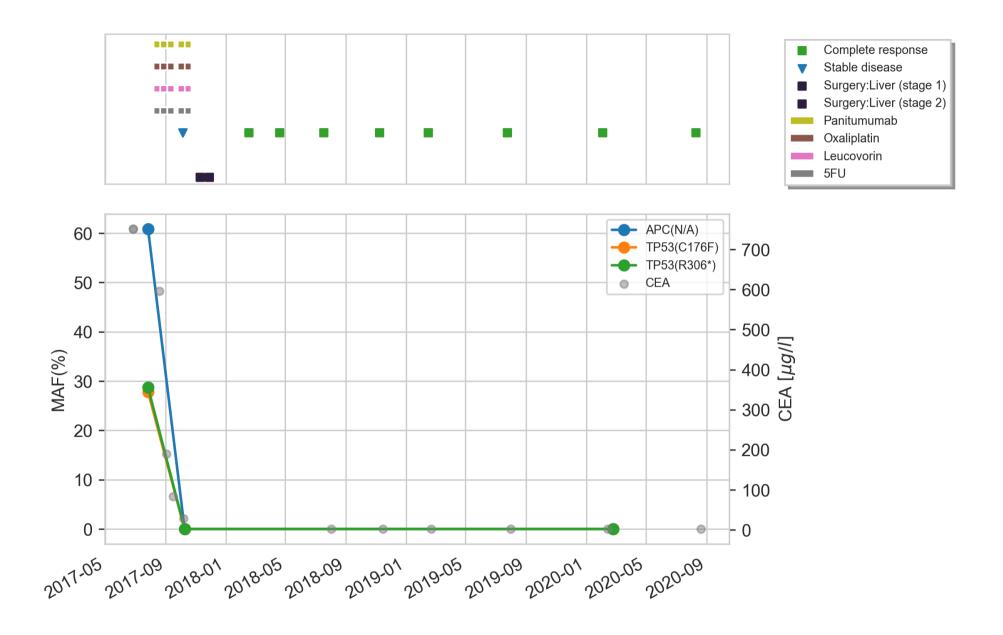
Patient 176

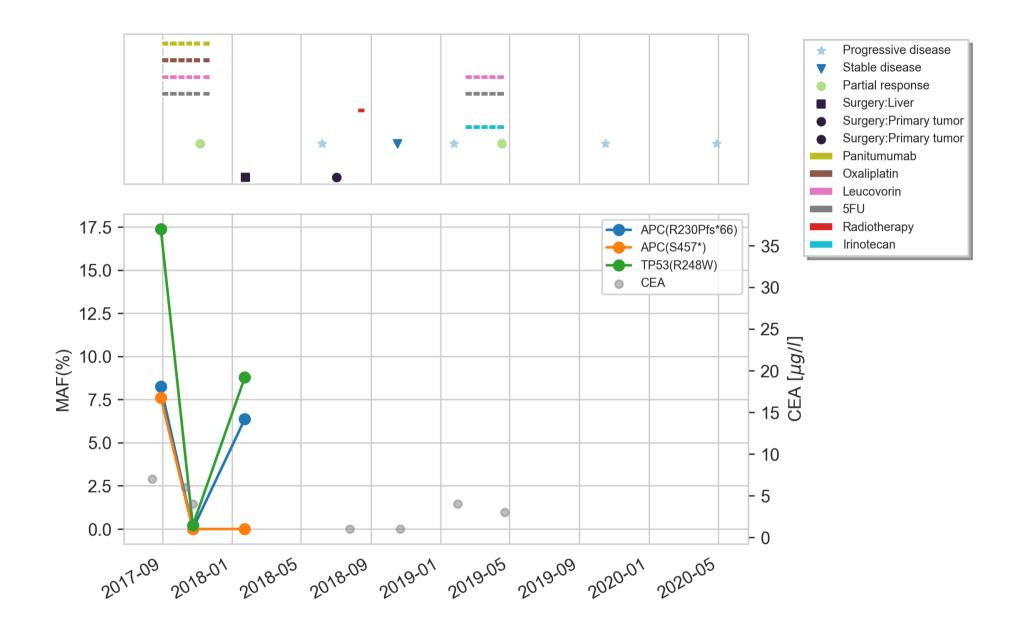




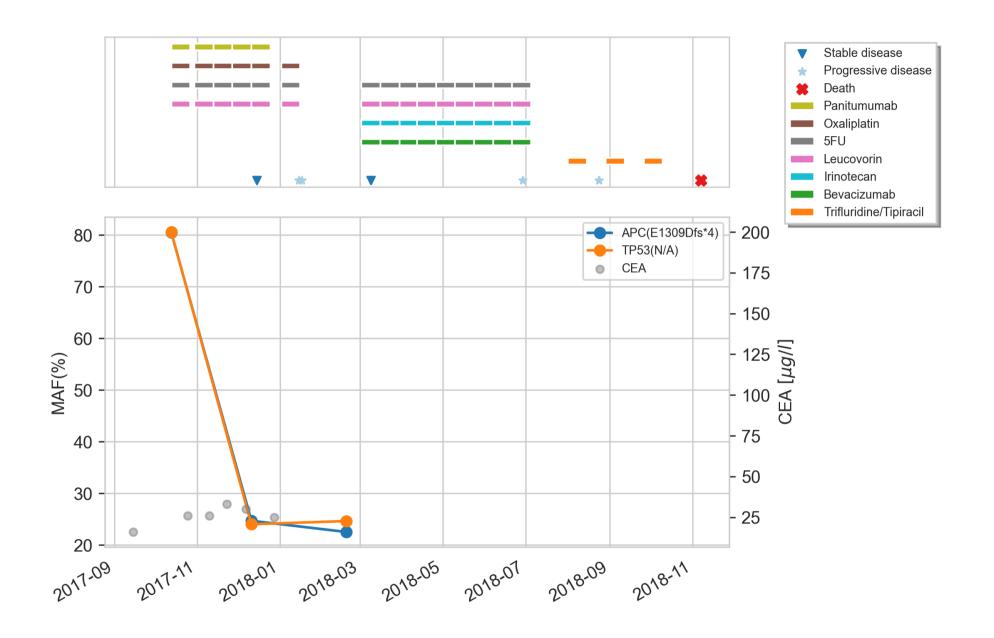


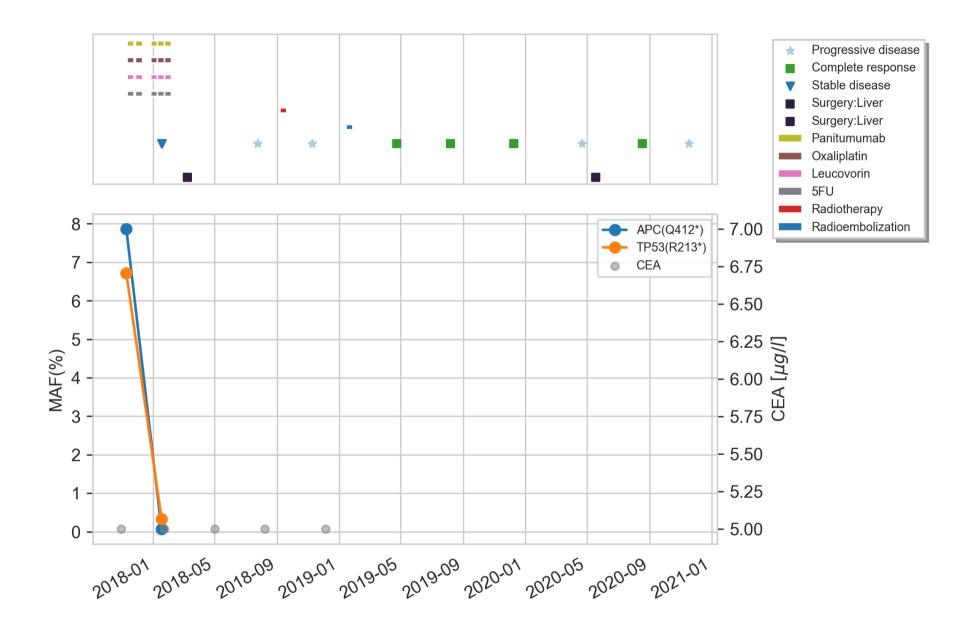






Patient 210





Patient 214

